

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/580,458A
Source: IFWO
Date Processed by STIC: 8/22/06

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 08/22/2006

PATENT APPLICATION: US/10/580,458A

TIME: 10:54:29

Input Set : A:\65691-445CorrSeqList.txt

Output Set: N:\CRF4\08222006\J580458A.raw

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3 <110> APPLICANT: Assistance Publique - Hopitaux de Paris (AH-HP)
4   Institut National de la Sante et de la Recherche Medicale
5   (INSERM)
6   Institut Gustave Roussy (IGR)
7   Universite de Versailles - Saint-Quentin-en-Yvelines
8   Universite Paris-Sud
9   VAINCHENKER, William
10  UGO, Valerie
11  JAMES, Chloe
12  LE COUEDIC, Jean-Pierre
13  CASADEVALL, Nicole
15 <120> TITLE OF INVENTION: Identification of a JAK2 mutation involved in Vaquez
16   Polyglobulia
18 <130> FILE REFERENCE: D 22707
C--> 20 <140> CURRENT APPLICATION NUMBER: US/10/580,458A
C--> 21 <141> CURRENT FILING DATE: 2006-05-24
23 <160> NUMBER OF SEQ ID NOS: 31
25 <170> SOFTWARE: PatentIn version 3.3
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 1132
29 <212> TYPE: PRT
30 <213> ORGANISM: homo sapiens
33 <220> FEATURE:
34 <223> OTHER INFORMATION: variant JAK2 V617F
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44 Lys Gln Ile Asp Pro Val Leu Gln Val Tyr Leu Tyr His Ser Leu Gly
45           35           40           45
47 Lys Ser Glu Ala Asp Tyr Leu Thr Phe Pro Ser Gly Glu Tyr Val Ala
48           50           55           60
50 Glu Glu Ile Cys Ile Ala Ala Ser Lys Ala Cys Gly Ile Thr Pro Val
51 65           70           75           80
53 Tyr His Asn Met Phe Ala Leu Met Ser Glu Thr Glu Arg Ile Trp Tyr
54           85           90           95
56 Pro Pro Asn His Val Phe His Ile Asp Glu Ser Thr Arg His Asn Val
57           100          105          110
59 Leu Tyr Arg Ile Arg Phe Tyr Phe Pro Arg Trp Tyr Cys Ser Gly Ser
60           115          120          125
62 Asn Arg Ala Tyr Arg His Gly Ile Ser Arg Gly Ala Glu Ala Pro Leu
63           130          135          140

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65 Leu Asp Asp Phe Val Met Ser Tyr Leu Phe Ala Gln Trp Arg His Asp
66 145 150 155 160
68 Phe Val His Gly Trp Ile Lys Val Pro Val Thr His Glu Thr Gln Glu
69 165 170 175
71 Glu Cys Leu Gly Met Ala Val Leu Asp Met Met Arg Ile Ala Lys Glu
72 180 185 190
74 Asn Asp Gln Thr Pro Leu Ala Ile Tyr Asn Ser Ile Ser Tyr Lys Thr
75 195 200 205
77 Phe Leu Pro Lys Cys Ile Arg Ala Lys Ile Gln Asp Tyr His Ile Leu
78 210 215 220
80 Thr Arg Lys Arg Ile Arg Tyr Arg Phe Arg Arg Phe Ile Gln Gln Phe
81 225 230 235 240
83 Ser Gln Cys Lys Ala Thr Ala Arg Asn Leu Lys Leu Lys Tyr Leu Ile
84 245 250 255
86 Asn Leu Glu Thr Leu Gln Ser Ala Phe Tyr Thr Glu Lys Phe Glu Val
87 260 265 270
89 Lys Glu Pro Gly Ser Gly Pro Ser Gly Glu Glu Ile Phe Ala Thr Ile
90 275 280 285
92 Ile Ile Thr Gly Asn Gly Gly Ile Gln Trp Ser Arg Gly Lys His Lys
93 290 295 300
95 Glu Ser Glu Thr Leu Thr Glu Gln Asp Leu Gln Leu Tyr Cys Asp Phe
96 305 310 315 320
98 Pro Asn Ile Ile Asp Val Ser Ile Lys Gln Ala Asn Gln Glu Gly Ser
99 325 330 335
101 Asn Glu Ser Arg Val Val Thr Ile His Lys Gln Asp Gly Lys Asn Leu
102 340 345 350
104 Glu Ile Glu Leu Ser Ser Leu Arg Glu Ala Leu Ser Phe Val Ser Leu
105 355 360 365
107 Ile Asp Gly Tyr Tyr Arg Leu Thr Ala Asp Ala His His Tyr Leu Cys
108 370 375 380
110 Lys Glu Val Ala Pro Pro Ala Val Leu Glu Asn Ile Gln Ser Asn Cys
111 385 390 395 400
113 His Gly Pro Ile Ser Met Asp Phe Ala Ile Ser Lys Leu Lys Lys Ala
114 405 410 415
116 Gly Asn Gln Thr Gly Leu Tyr Val Leu Arg Cys Ser Pro Lys Asp Phe
117 420 425 430
119 Asn Lys Tyr Phe Leu Thr Phe Ala Val Glu Arg Glu Asn Val Ile Glu
120 435 440 445
122 Tyr Lys His Cys Leu Ile Thr Lys Asn Glu Asn Glu Glu Tyr Asn Leu
123 450 455 460
125 Ser Gly Thr Lys Lys Asn Phe Ser Ser Leu Lys Asp Leu Leu Asn Cys
126 465 470 475 480
128 Tyr Gln Met Glu Thr Val Arg Ser Asp Asn Ile Ile Phe Gln Phe Thr
129 485 490 495
131 Lys Cys Cys Pro Pro Lys Pro Lys Asp Lys Ser Asn Leu Leu Val Phe
132 500 505 510
134 Arg Thr Asn Gly Val Ser Asp Val Pro Thr Ser Pro Thr Leu Gln Arg
135 515 520 525
137 Pro Thr His Met Asn Gln Met Val Phe His Lys Ile Arg Asn Glu Asp

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213      930      935      940
215 Lys Glu Arg Ile Asp His Ile Lys Leu Leu Gln Tyr Thr Ser Gln Ile
216 945      950      955      960
218 Cys Lys Gly Met Glu Tyr Leu Gly Thr Lys Arg Tyr Ile His Arg Asp
219      965      970      975
221 Leu Ala Thr Arg Asn Ile Leu Val Glu Asn Glu Asn Arg Val Lys Ile
222      980      985      990
224 Gly Asp Phe Gly Leu Thr Lys Val Leu Pro Gln Asp Lys Glu Tyr Tyr
225      995      1000      1005
227 Lys Val Lys Glu Pro Gly Glu Ser Pro Ile Phe Trp Tyr Ala Pro
228      1010      1015      1020
230 Glu Ser Leu Thr Glu Ser Lys Phe Ser Val Ala Ser Asp Val Trp
231      1025      1030      1035
233 Ser Phe Gly Val Val Leu Tyr Glu Leu Phe Thr Tyr Ile Glu Lys
234      1040      1045      1050
236 Ser Lys Ser Pro Pro Ala Glu Phe Met Arg Met Ile Gly Asn Asp
237      1055      1060      1065
239 Lys Gln Gly Gln Met Ile Val Phe His Leu Ile Glu Leu Leu Lys
240      1070      1075      1080
242 Asn Asn Gly Arg Leu Pro Arg Pro Asp Gly Cys Pro Asp Glu Ile
243      1085      1090      1095
245 Tyr Met Ile Met Thr Glu Cys Trp Asn Asn Asn Val Asn Gln Arg
246      1100      1105      1110
248 Pro Ser Phe Arg Asp Leu Ala Leu Arg Val Asp Gln Ile Arg Asp
249      1115      1120      1125
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252      1130
254 <210> SEQ ID NO: 2
255 <211> LENGTH: 5097
256 <212> TYPE: DNA
257 <213> ORGANISM: homo sapiens
260 <220> FEATURE:
261 <223> OTHER INFORMATION: G1849T mutation in jak2 gene
263 <400> SEQUENCE: 2
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268 tcggcttctc ggccggtcgg gccctcggc ccgggcttgc ggcgcgcgtc ggggctgagg      180
270 gctgctgcgg cgcagggaga ggcctggtcc tcgctgccga gggatgtgag tgggagctga      240
272 gccacactg gagggccccc gagggcccag cctggaggtc gttcagagcc gtgcccggcc      300
274 cggggcttcg cagaccttga cccgccgggt aggagccgcc cctgcgggct cgagggcgcg      360
276 ctctggtcgc ccgatctgtg tagccggttt cagaagcagg caacaggaac aagatgtgaa      420
278 ctgtttctct tctgcagaaa aagaggctct tctcctcct cccgcgacgg caaatgttct      480
280 gaaaaagact ctgcatggga atggcctgcc ttacgatgac agaaatggag ggaacatcca      540
282 cctcttctat atatcagaat ggtgatattt ctggaaatgc caattctatg aagcaaatag      600
284 atccagttct tcaggtgtat ctttaccatt cccttgggaa atctgaggca gattatctga      660
286 cctttccatc tggggagtat gttgcagaag aaatctgtat tgctgcttct aaagcttgtg      720
288 gtatcacacc tgtgtatcat aatatgtttg ctttaatgag tgaaacagaa aggatctggt      780
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296	agtggcggca	tgattttgtg	cacggatgga	taaaagtacc	tgtgactcat	gaaacacagg	1020
298	aagaatgtct	tgggatggca	gtgttagata	tgatgagaat	agccaaagaa	aacgatcaaa	1080
300	ccccactggc	catctataac	tctatcagct	acaagacatt	cttaccaaaa	tgtattcgag	1140
302	caaagatcca	agactatcat	attttgacaa	ggaagcgaat	aaggtacaga	tttcgcagat	1200
304	ttattcagca	attcagccaa	tgcaaagcca	ctgccagaaa	cttgaaactt	aagtatctta	1260
306	taaatctgga	aactctgcag	tctgccttct	acacagagaa	atttgaagta	aaagaacctg	1320
308	gaagtgggtc	ttcaggtgag	gagatttttg	caaccattat	aataactgga	aacggtggaa	1380
310	ttcagtggtc	aagagggaaa	cataaagaaa	gtgagacact	gacagaacag	gatttacagt	1440
312	tatattgcca	ttttccta	attattgatg	tcagtattaa	gcaagcaa	caagagggtt	1500
314	caaataagaa	ccgagttgta	actatccata	agcaagatgg	taaaaatctg	gaaattgaac	1560
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318	ctgcagatgc	acatcattac	ctctgtaaa	aagtagcacc	tccagccgtg	cttgaaaata	1680
320	tacaaagcaa	ctgtcatggc	ccaatttcga	tggattttgc	cattagtaaa	ctgaagaaag	1740
322	caggtaatca	gactggactg	tatgtacttc	gatgcagtcc	taaggacttt	aataaatatt	1800
324	ttttgacttt	tgctgtcgag	cgagaaaatg	tcattgaata	taaacactgt	ttgattacaa	1860
326	aaaatgagaa	tgaagagtac	aacctcagtg	ggacaaagaa	gaacttcagc	agtcttaaa	1920
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338	cagaagttct	tttaaaagtt	ctggataaag	cacacagaaa	ctattcagag	tctttctttg	2280
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360	cttctttcag	agccatcata	cgagatctta	acagtttgtt	tactccagat	tatgaactat	2940
362	taacagaaaa	tgacatgtta	ccaaatatga	ggataggtgc	cctaggggtt	tctgggtgct	3000
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384	atgtttggag	ctttggagtg	gttctgtatg	aacttttcac	atacattgag	aagagtaaaa	3660
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/580,458A

DATE: 08/22/2006
TIME: 10:54:31

Input Set : A:\65691-445CorrSeqList.txt
Output Set: N:\CRF4\08222006\J580458A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; N Pos. 20,21

Seq#:14; N Pos. 20,21

VERIFICATION SUMMARY

DATE: 08/22/2006

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Input Set : A:\65691-445CorrSeqList.txt

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L:20 M:270 C: Current Application Number differs, Replaced Current Application Number

L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0

L:622 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0